

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/561,144
Source: IFWP
Date Processed by STIC: 12/30/05

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 12/30/2005

PATENT APPLICATION: US/10/561,144

TIME: 09:39:19

Input Set : A:\1431sequence_listing.txt

Output Set: N:\CRF4\12302005\J561144.raw

3 <110> APPLICANT: TAKEDA CHEMICAL INDUSTRIES, LTD.
 4 SHIMOMURA, Ichirou
 6 <120> TITLE OF INVENTION: Novel Protein
 8 <130> FILE REFERENCE: 09653
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/561,144
 C--> 10 <141> CURRENT FILING DATE: 2005-12-16
 10 <150> PRIOR APPLICATION NUMBER: JP 2003-171188
 11 <151> PRIOR FILING DATE: 2003-06-16
 13 <150> PRIOR APPLICATION NUMBER: JP 2003-391047
 14 <151> PRIOR FILING DATE: 2003-11-20
 16 <150> PRIOR APPLICATION NUMBER: JP 2004-23557
 17 <151> PRIOR FILING DATE: 2004-01-30
 19 <150> PRIOR APPLICATION NUMBER: JP 2004-30988
 20 <151> PRIOR FILING DATE: 2004-02-06
 22 <160> NUMBER OF SEQ ID NOS: 25
 24 <170> SOFTWARE: PatentIn version 3.2
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 402
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapiens
 32 <220> FEATURE:
 33 <221> NAME/KEY: CDS
 34 <222> LOCATION: (1)..(399)
 36 <220> FEATURE:
 37 <221> NAME/KEY: sig_peptide
 38 <222> LOCATION: (1)..(87)
 40 <220> FEATURE:
 41 <221> NAME/KEY: mat_peptide
 42 <222> LOCATION: (88)..(399)
 44 <400> SEQUENCE: 1
 45 atg ctg gac tgg aga ttg gca agt gca cat ttc atc ctg gct gtg aca 48
 46 Met Leu Asp Trp Arg Leu Ala Ser Ala His Phe Ile Leu Ala Val Thr
 47 -25 -20 -15
 49 ctg aca ctg tgg agc tca gga aaa gtc ctc tca gta gat gta aca aca 96
 50 Leu Thr Leu Trp Ser Ser Gly Lys Val Leu Ser Val Asp Val Thr Thr
 51 -10 -5 -1 1
 53 aca gag gcc ttt gat tct gga gtc ata gat gtg cag tca aca ccc aca 144
 54 Thr Glu Ala Phe Asp Ser Gly Val Ile Asp Val Gln Ser Thr Pro Thr
 55 5 10 15
 57 gtc agg gaa gag aaa tca gcc act gac ctg aca gca aaa ctc ttg ctt 192
 58 Val Arg Glu Glu Lys Ser Ala Thr Asp Leu Thr Ala Lys Leu Leu Leu
 59 20 25 30 35
 61 ctt gat gaa ttg gtg tcc cta gaa aat gat gtg att gag aca aag aag 240

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62 Leu Asp Glu Leu Val Ser Leu Glu Asn Asp Val Ile Glu Thr Lys Lys
63          40          45          50
65 aaa agg agt ttc tct ggt ttt ggg tct ccc ctt gac aga ctc tca gct      288
66 Lys Arg Ser Phe Ser Gly Phe Gly Ser Pro Leu Asp Arg Leu Ser Ala
67          55          60          65
69 ggc tct gta gat cac aaa ggt aaa cag agg aaa gta gta gat cat cca      336
70 Gly Ser Val Asp His Lys Gly Lys Gln Arg Lys Val Val Asp His Pro
71          70          75          80
73 aaa agg cga ttt ggt atc ccc atg gat cgg att ggt aga aac cgg ctt      384
74 Lys Arg Arg Phe Gly Ile Pro Met Asp Arg Ile Gly Arg Asn Arg Leu
75          85          90          95
77 tca aat tcc aga ggc taa      402
78 Ser Asn Ser Arg Gly
79 100
82 <210> SEQ ID NO: 2
83 <211> LENGTH: 133
84 <212> TYPE: PRT
85 <213> ORGANISM: Homo sapiens
87 <400> SEQUENCE: 2
89 Met Leu Asp Trp Arg Leu Ala Ser Ala His Phe Ile Leu Ala Val Thr
90          -25          -20          -15
93 Leu Thr Leu Trp Ser Ser Gly Lys Val Leu Ser Val Asp Val Thr Thr
94          -10          -5          -1 1
97 Thr Glu Ala Phe Asp Ser Gly Val Ile Asp Val Gln Ser Thr Pro Thr
98          5          10          15
101 Val Arg Glu Glu Lys Ser Ala Thr Asp Leu Thr Ala Lys Leu Leu Leu
102 20          25          30          35
105 Leu Asp Glu Leu Val Ser Leu Glu Asn Asp Val Ile Glu Thr Lys Lys
106          40          45          50
109 Lys Arg Ser Phe Ser Gly Phe Gly Ser Pro Leu Asp Arg Leu Ser Ala
110          55          60          65
113 Gly Ser Val Asp His Lys Gly Lys Gln Arg Lys Val Val Asp His Pro
114          70          75          80
117 Lys Arg Arg Phe Gly Ile Pro Met Asp Arg Ile Gly Arg Asn Arg Leu
118          85          90          95
121 Ser Asn Ser Arg Gly
122 100
125 <210> SEQ ID NO: 3
126 <211> LENGTH: 393
127 <212> TYPE: DNA
128 <213> ORGANISM: Mus musculus
131 <220> FEATURE:
132 <221> NAME/KEY: CDS
133 <222> LOCATION: (1)..(390)
135 <220> FEATURE:
136 <221> NAME/KEY: sig_peptide
137 <222> LOCATION: (1)..(87)
139 <220> FEATURE:
140 <221> NAME/KEY: mat_peptide

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141 <222> LOCATION: (88)..(390)
143 <400> SEQUENCE: 3
144 atg ctg gac tgg aga ttg gca agt aca cac ttc atc ctg gct atg att      48
145 Met Leu Asp Trp Arg Leu Ala Ser Thr His Phe Ile Leu Ala Met Ile
146          -25          -20          -15
148 gtg atg ctg tgg ggc tca gga aag gca ttc tct gtg gac tta gca tca      96
149 Val Met Leu Trp Gly Ser Gly Lys Ala Phe Ser Val Asp Leu Ala Ser
150          -10          -5          -1  1
152 cag gag ttt gga aca gca agc ttg cag tct cca ccc aca gcc aga gaa      144
153 Gln Glu Phe Gly Thr Ala Ser Leu Gln Ser Pro Pro Thr Ala Arg Glu
154    5          10          15
156 gag aag tca gcc act gag ctt tcg gct aag ctc ctg cgt ctt gat gat      192
157 Glu Lys Ser Ala Thr Glu Leu Ser Ala Lys Leu Leu Arg Leu Asp Asp
158 20          25          30          35
160 ctg gtg tcc tta gag aat gac gta ttt gag acc aag aaa aag aga agc      240
161 Leu Val Ser Leu Glu Asn Asp Val Phe Glu Thr Lys Lys Lys Arg Ser
162          40          45          50
164 ttc tct ggc ttt ggg tct ccc ctt gac aga ctc tca gct ggg tct gta      288
165 Phe Ser Gly Phe Gly Ser Pro Leu Asp Arg Leu Ser Ala Gly Ser Val
166          55          60          65
168 gag cat aga ggg aaa caa agg aaa gca gta gat cat tca aaa aag cgg      336
169 Glu His Arg Gly Lys Gln Arg Lys Ala Val Asp His Ser Lys Lys Arg
170          70          75          80
172 ttt ggt att ccc atg gat cgg att ggt aga aac cgg ctc tcc agt tcc      384
173 Phe Gly Ile Pro Met Asp Arg Ile Gly Arg Asn Arg Leu Ser Ser Ser
174    85          90          95
176 aga ggc tga      393
177 Arg Gly
178 100
181 <210> SEQ ID NO: 4
182 <211> LENGTH: 130
183 <212> TYPE: PRT
184 <213> ORGANISM: Mus musculus
186 <400> SEQUENCE: 4
188 Met Leu Asp Trp Arg Leu Ala Ser Thr His Phe Ile Leu Ala Met Ile
189          -25          -20          -15
192 Val Met Leu Trp Gly Ser Gly Lys Ala Phe Ser Val Asp Leu Ala Ser
193          -10          -5          -1  1
196 Gln Glu Phe Gly Thr Ala Ser Leu Gln Ser Pro Pro Thr Ala Arg Glu
197    5          10          15
200 Glu Lys Ser Ala Thr Glu Leu Ser Ala Lys Leu Leu Arg Leu Asp Asp
201 20          25          30          35
204 Leu Val Ser Leu Glu Asn Asp Val Phe Glu Thr Lys Lys Lys Arg Ser
205          40          45          50
208 Phe Ser Gly Phe Gly Ser Pro Leu Asp Arg Leu Ser Ala Gly Ser Val
209          55          60          65
212 Glu His Arg Gly Lys Gln Arg Lys Ala Val Asp His Ser Lys Lys Arg
213          70          75          80
216 Phe Gly Ile Pro Met Asp Arg Ile Gly Arg Asn Arg Leu Ser Ser Ser

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217      85              90              95
220 Arg Gly
221 100
224 <210> SEQ ID NO: 5
225 <211> LENGTH: 19
226 <212> TYPE: DNA
227 <213> ORGANISM: Artificial Sequence
229 <220> FEATURE:
230 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
cDNA
231      derived from mouse soleus muscle.
233 <400> SEQUENCE: 5
234 gggggtggac catcctcta                      19
237 <210> SEQ ID NO: 6
238 <211> LENGTH: 20
239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial Sequence
242 <220> FEATURE:
243 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
cDNA
244      derived from mouse soleus muscle.
246 <400> SEQUENCE: 6
247 cgcgcagctg taaacggtag                      20
250 <210> SEQ ID NO: 7
251 <211> LENGTH: 249
252 <212> TYPE: DNA
253 <213> ORGANISM: Mus musculus
255 <400> SEQUENCE: 7
256 aggactctaa agttaggagc tctgacttct cacaagatgc tggactggag attggcaagt      60
258 acacacttca tcctggctat gattgtgatg ctgtggggct caggaaaggc attctctgtg      120
260 gacttagcat cacaggagtt tggaacagca agcttgacgt ctccaccac agccagagaa      180
262 gagaagtcag ccaactgagct ttcggctaag ctccctgcgtc ttgatgatct ggtgtcctta      240
264 gagaatgac                                249
267 <210> SEQ ID NO: 8
268 <211> LENGTH: 30
269 <212> TYPE: DNA
270 <213> ORGANISM: Artificial Sequence
272 <220> FEATURE:
273 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
full
274      length mouse muscudin cDNA.
276 <400> SEQUENCE: 8
277 tcctgagccc cacagcatca caatcatagc                      30
280 <210> SEQ ID NO: 9
281 <211> LENGTH: 19
282 <212> TYPE: DNA
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
full
287      length mouse muscudin cDNA.
289 <400> SEQUENCE: 9
290 ccttgacaga ctctcagct                                19

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Input Set : A:\1431sequence_listing.txt

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293 <210> SEQ ID NO: 10
294 <211> LENGTH: 1154
295 <212> TYPE: DNA
296 <213> ORGANISM: Mus musculus
298 <400> SEQUENCE: 10
299 gagagagaga gagagagaga gagagagaga gagagttggt gaaatgttcc gctgaaaatc      60
301 tgtggaactg atgtaagaga aagcaacgac aggggttgga gtaagtggag tagaactgag      120
303 actataaaaa cacagaaaga aactcgcatc agggctaagt ttgggataag ctgcaggcag      180
305 gactctaaag ttaggagctc tgactttctca caagatgctg gactggagat tggcaagtac      240
307 acacttcatac ctggctatga ttgtgatgct gtggggctca ggaaaggcat tctctgtgga      300
309 cttagcatca caggagtttg gaacagcaag cttgcagtct ccaccacag ccagagaaga      360
311 gaagtcagcc actgagcttt cggctaagct cctgcgtctt gatgatctgg tgccttaga      420
313 gaatgacgta tttagacca agaaaaagag aagcttctct ggctttgggt ctccccttga      480
315 cagactctca gctgggtctg tagagcatag agggaaacaa aggaaagcag tagatcattc      540
317 aaaaaagcgg tttggtattc ccattgacg gattggtaga aaccggctct ccagttccag      600
319 agggctgatg attcttattg tgcgacttac ttgtgtgaga tggcacagaa ctatagaaga      660
321 cacttcagtg aagttcacta ccccttttgt caaggaattg gcctttcgca aaccttccca      720
323 aagcttgatc ctccccagac catcacgtca tagtgttgct gtggttttag ttgagttgtg      780
325 cagatcattt cagtgcattg atatctctga aagtattttt caatgattcc caaattgtaa      840
327 cgtggccctt gaacctactt tttttaaaca gcagaccaat ataatgcatt ctcttgccat      900
329 taatattttt acatttcagt taatcaatgt gctttctaga aacctagtgt ttgaagatct      960
331 gatgatctaa agaaatcaga aatgagcaca tggtgattta tataggtttc tttagttttt     1020
333 ctgagggttg tcgaattggt gtaaacttca acttcaagct tagaaaaaag acattacatg     1080
335 agtgtttgct tcaactgtgt cagaaggcaa ataaattttg agaaaccaa aaaaaaaaaa     1140
337 aaaaaaaaaa aaaa                                         1154
340 <210> SEQ ID NO: 11
341 <211> LENGTH: 28
342 <212> TYPE: DNA
343 <213> ORGANISM: Artificial Sequence
345 <220> FEATURE:
346 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
full
347         length human muscudin cDNA.
349 <400> SEQUENCE: 11
350 gactgtgggt gttgactgca catctatg                                         28
353 <210> SEQ ID NO: 12
354 <211> LENGTH: 28
355 <212> TYPE: DNA
356 <213> ORGANISM: Artificial Sequence
358 <220> FEATURE:
359 <223> OTHER INFORMATION: Oligonucleotide designed to act as primer for amplifying
full
360         length human muscudin cDNA.
362 <400> SEQUENCE: 12
363 ctgatctttc acgagatgct ggactgga                                         28
366 <210> SEQ ID NO: 13
367 <211> LENGTH: 1655
368 <212> TYPE: DNA
369 <213> ORGANISM: Homo sapiens
372 <220> FEATURE:
373 <221> NAME/KEY: misc_feature

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 12/30/2005
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; N Pos. 763,965,1032,1033,1043,1075,1080,1113,1138,1235,1238,1265

Seq#:13; N Pos. 1277,1280,1291,1301,1328,1347,1366,1454,1553,1580,1625

VERIFICATION SUMMARY

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:512 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:720
L:520 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:960
L:522 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1020
L:524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1080
L:528 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1200
L:530 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1260
L:532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1320
L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1440
L:538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1500
L:540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1560
L:542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:1620